

# Interleukin-2 Inhibits Germinal Center Formation by Limiting T Follicular Helper Cell Differentiation

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## SUMMARY

T follicular helper (Tfh) cells promote T cell-dependent humoral immune responses by providing T cell help to B cells and by promoting germinal center (GC) formation and long-lived antibody responses. However, the cellular and molecular mechanisms that control Tfh cell differentiation *in vivo* are incompletely understood. Here we show that interleukin-2 (IL-2) administration impaired influenza-specific GCs, long-lived IgG responses, and Tfh cells. IL-2 did not directly inhibit GC formation, but instead suppressed the differentiation of Tfh cells, thereby hindering the maintenance of influenza-specific GC B cells. Our data demonstrate that IL-2 is a critical factor that regulates successful Tfh and B cell responses *in vivo* and regulates Tfh cell development.

## INTRODUCTION

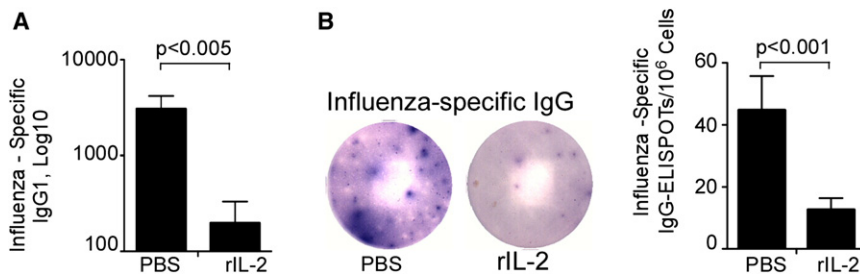
The generation of long-lived plasma cells that secrete high-affinity antibodies is critical for the establishment of protective immunity to influenza and other pathogens. The selection of long-lived B cells with high-affinity BCRs is dependent on events that occur in the germinal center (GC) (Allen et al., 2007; Cyster, 2010; Kelsoe, 1996). GCs are structures found in the B cell follicles of reactive lymphoid tissues and support intense B cell clonal expansion, somatic hypermutation of immunoglobulin genes, affinity maturation of the B cell response, and ultimately the differentiation of both memory B cells and long-lived plasma cells (Allen et al., 2007; Cyster, 2010; Vinuesa et al., 2010). Given that the goal of most vaccination strategies is the generation of long-lived antibody responses (Cox et al., 2004), it is essential that we understand the factors that regulate GC formation and maintenance.

GCs are dependent on help provided by CD4<sup>+</sup> T follicular helper (Tfh) cells—a recently defined CD4<sup>+</sup> T cell subset with a unique ability to help B cell responses (Crotty, 2011; Yu and Vinuesa, 2010). B cells and Tfh cells provide each other with reciprocal differentiation and survival signals (King, 2011). Tfh cells provide CD40 ligand (Goodnow et al., 2010; Han et al., 1995) and IL-21 to B cells (Bryant et al., 2007; Linterman et al., 2010; Vogelzang et al., 2008) and B cells present antigen and provide ICOS ligand to Tfh cells (Choi et al., 2011; Deenick

et al., 2011; Johnston et al., 2009; Nurieva et al., 2009). As a result, all three of these molecules are essential for germinal center formation and long-lived antibody responses (Goodnow et al., 2010). Importantly, the differentiation of Tfh cells depends on the expression of the transcription factor Bcl6, which represses the expression of other lineage-specific transcription factors and promotes the expression of CXCR5 (Johnston et al., 2009; Nurieva et al., 2009; Yu et al., 2009). In turn, CXCR5 facilitates homing to the B cell follicle and germinal center (Hardtke et al., 2005; Haynes et al., 2007). As a result, Bcl6-deficient CD4<sup>+</sup> T cells do not properly express CXCR5, fail to migrate into B cell follicles, and do not support GC reactions. Thus, Bcl6 is considered the master regulator of Tfh cell differentiation (Johnston et al., 2009; Nurieva et al., 2009; Yu et al., 2009). However, the molecular mechanisms that regulate Bcl6 expression and the differentiation of Tfh cells *in vivo* are incompletely defined (Crotty, 2011; Yu and Vinuesa, 2010).

Recent results demonstrate that CD4<sup>+</sup> T cells responding to LCMV can be separated based on the expression of the interleukin-2 receptor  $\alpha$  (CD25) into CD25<sup>hi</sup> and CD25<sup>lo</sup> subsets (Choi et al., 2011). Importantly, CD25<sup>hi</sup> T cells do not express Bcl6, suggesting a possible role for IL-2 in the segregation of effector and Tfh cells. IL-2 was originally defined as an essential T cell growth factor that promotes expansion of effector T cells and facilitates CD8<sup>+</sup> memory T cell programming (Blattman et al., 2003; Malek, 2008; Malek and Castro, 2010; Williams et al., 2006). However, IL-2 signaling also controls the contraction of T cell responses by enhancing susceptibility of responding T cells to activation-induced cell death (Lenardo, 1991). IL-2 signaling through CD25 enhances CD25 expression and sensitivity to IL-2 and also controls the expression of key transcription factors, such as eomesodermin (Eomes), Blimp1, and Bcl6, in CD8 T cells (Pipkin et al., 2010).

The generation and maintenance of FoxP3-expressing CD4<sup>+</sup> regulatory T (Treg) cells are also dependent on IL-2 (Almeida et al., 2002; de la Rosa et al., 2004; Malek and Castro, 2010; Malek et al., 2002). Treg cells suppress the activation of autoreactive T cells, prevent the development autoimmune disease (Campbell and Koch, 2011; Malek et al., 2002), and play an important role in the control of B cell responses (Alexander et al., 2011; Chung et al., 2011; Lim et al., 2004, 2005; Linterman et al., 2011). Given the essential role for IL-2 in the maintenance of Treg cells, some investigators are using IL-2 therapy to treat autoimmune disease in mouse models (Grinberg-Bleyer et al., 2010; Humrich et al., 2010). In humans, IL-2 therapy has been used to enhance antitumor immunity (Kovacs et al., 2001;



**Figure 1. IL-2 Inhibits B Cell Responses to Influenza**

B6 mice were infected with PR8 and treated with 30,000 U of human rIL-2 or PBS daily starting 3 days after infection.

(A) Serum was obtained on day 21 after infection and titers of influenza-specific IgG1 were determined by ELISA.

(B) Influenza-specific IgG-secreting cells in the BM were enumerated by ELISPOT on day 21 after infection.

Data are representative of two independent experiments (mean  $\pm$  SD of five mice per group).

Rosenberg et al., 1985a, 1985b), particularly in combination with B cell depletion in the treatment of non-Hodgkin lymphoma (Friedberg et al., 2002; Gillies et al., 2005; Gluck et al., 2004; Lopes de Menezes et al., 2007). Thus, IL-2 may be used to treat a variety of diseases.

Although IL-2 can modulate effector and Treg cell function by a variety of mechanisms, the role of IL-2 in the control of B cell and Tfh cell responses remain unclear. Here we tested the role of IL-2 in the GC B cell response to influenza. We found that IL-2 impaired the influenza-specific GC response, reduced the influenza-specific IgG response, and prevented the accumulation of influenza-specific Tfh cells. We demonstrated that IL-2 did not directly inhibit GCs, but instead directly suppressed Tfh cell responses. We found that CD25 signaling triggered by IL-2 negatively modulated Bcl6 expression and Tfh cell development. These data demonstrate that the physiological availability of IL-2 is a critical factor that regulates successful Tfh and B cell responses in vivo.

## RESULTS

### IL-2 Treatment Impairs GC B Cell Formation and Antibody Responses after Influenza Infection

To determine the role of IL-2 in the B cell response to influenza, we infected C57BL/6 (B6) mice with influenza virus A/PR8/34 (PR8) and treated infected mice with 30,000 U of human recombinant IL-2 (rIL-2) or PBS twice a day starting 3 days after infection. On day 21 postinfection, we collected sera from rIL-2-treated and control mice and determined the titers of influenza-specific IgG by ELISA (Figure 1A). We observed a substantial reduction in influenza-specific IgG1 titers in the rIL-2-treated group compared to control mice. To determine whether rIL-2 treatment specifically blocked the generation of influenza-specific IgG-secreting plasma cells that home to the bone marrow (BM) and provide long-lasting immunity to influenza, we enumerated influenza-specific IgG-secreting plasma cells in the BM on day 21 postinfection by ELISPOT. In agreement with the antibody titers, we found that the number of IgG-secreting influenza-specific plasma cells in the bone marrow was reduced in the BM of rIL-2-treated mice relative to control mice (Figure 1B). Thus, our results indicate that rIL-2 treatment compromised influenza-specific humoral immune responses.

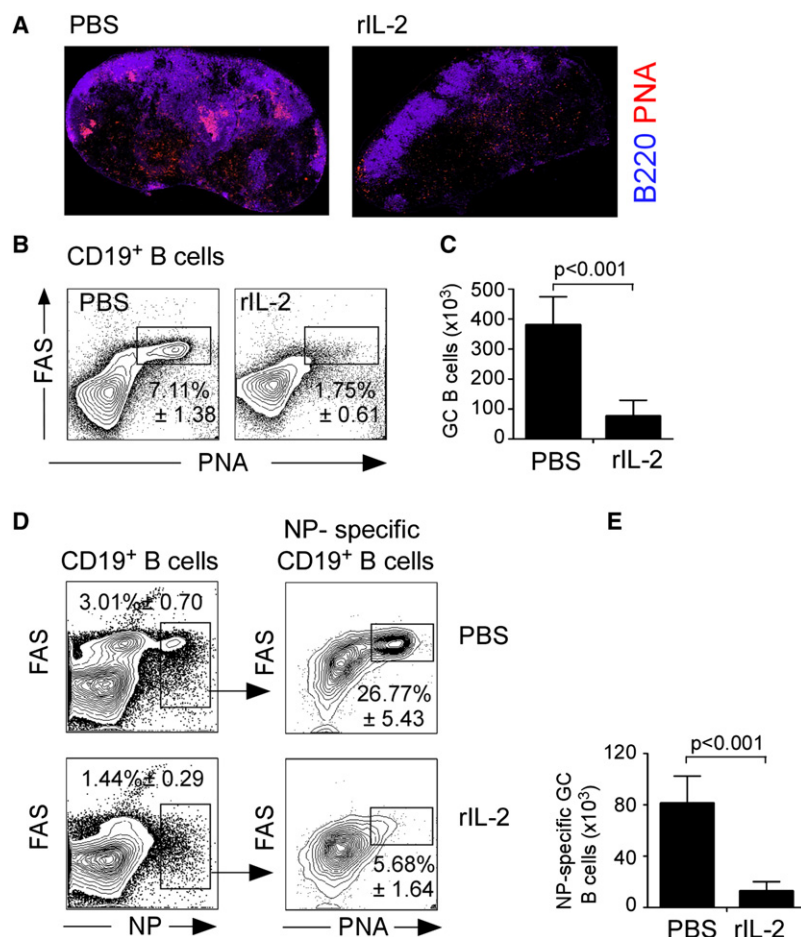
Given that GC formation is required for the differentiation of long-lived plasma cells (Johnston et al., 2009; Nurieva et al., 2009; Vinuesa et al., 2010; Yu et al., 2009), we hypothesized

that rIL-2 treatment may inhibit the formation of GCs after influenza infection. To test this hypothesis, we infected B6 mice with influenza and treated them with either rIL-2 or PBS. On day 10 postinfection, we probed cryosections of mediastinal lymph nodes (mLNs) with fluorochrome-labeled PNA and anti-B220 to detect GCs. As expected, PNA<sup>+</sup> GC structures were easily detected in the B cell follicles of control mice. However, although B cell follicles were present in rIL-2-treated mice, PNA<sup>+</sup> GC structures were virtually undetectable after rIL-2 treatment (Figure 2A). Consistent with the histological results, the frequency (Figure 2B) and number (Figure 2C) of CD19<sup>+</sup> PNA<sup>hi</sup>FAS<sup>+</sup> GC B cells were reduced in rIL-2-treated mice.

To determine whether the IL-2 treatment impaired the development of influenza-specific GC B cell responses, we used fluorochrome-labeled recombinant influenza nucleoprotein (NP) tetramers to identify NP-specific PNA<sup>hi</sup>FAS<sup>+</sup> GC B cells. We found that rIL-2 treatment reduced the frequency of NP-specific PNA<sup>hi</sup>FAS<sup>+</sup> GC B cells by nearly 80% compared to control mice (Figure 2D). In a control experiment, we demonstrated that the NP B cell tetramer bound only B cells from influenza-infected mice and not B cells from mice infected with other pathogens (Figure S1 available online). Collectively, these data show that the influenza-specific GC B cell response was severely compromised in the presence of excess IL-2.

### IL-2 Indirectly Suppresses the GC B Cell Response to Influenza

To determine whether rIL-2 treatment acted directly on GC B cells, we generated 50:50 mixed BM chimeras by using B6.CD45.1 bone marrow (BM) and Cd25<sup>-/-</sup> BM (CD45.2). As expected, the cells that repopulated the chimeric mice were derived equally from both donors (Figure 3A). We subsequently infected the chimeric mice with influenza, administered either rIL-2 or PBS, and determined the frequency of PNA<sup>hi</sup>FAS<sup>+</sup> GC cells within the B6 and Cd25<sup>-/-</sup> CD19<sup>+</sup> B cells on day 10. We found that B6 (CD45.1) and Cd25<sup>-/-</sup> (CD45.2) B cells contained equal frequencies of GC B cells in both PBS-treated (Figure 3B) and IL-2-treated (Figure 3C) mice. Although the GC B cell response was diminished in the rIL-2-treated chimeras relative to that in the PBS-treated mice (Figures 3B and 3C), the ratio of B6 to Cd25<sup>-/-</sup> GC B cells was close to 1.0 in both groups (Figure 3D). We observed similar results when we examined NP-specific GC B cell response of B6 and Cd25<sup>-/-</sup> donors (Figures 3E–3G). These results suggest that IL-2 inhibited the development of GC B cells indirectly and not through a B cell-intrinsic mechanism.



**Figure 2. IL-2 Impairs GC B Cell Responses to Influenza**

(A) B6 mice were infected with PR8 and treated with 30,000 U of human recombinant rIL-2 or PBS twice a day starting 3 days after infection. mLNs were obtained on day 10 after infection and cryosections were stained with anti-B220 (blue) and PNA (red) and analyzed by fluorescent microscopy.

(B and C) Cells from the mLNs of mice treated with IL-2 or PBS were obtained on day 10 and the percentage of CD19<sup>+</sup> B cells with a PNA<sup>+</sup>FAS<sup>+</sup> GC phenotype was determined (B), and the number of GC B cells was calculated (C). Data are representative of four independent experiments (mean  $\pm$  SD of five mice per group).

(D and E) Cells from mLNs were obtained on day 10 after infection and GC B cells were identified by flow cytometry with an NP tetramer. The percentage of NP-specific B cells with a PNA<sup>+</sup>FAS<sup>+</sup> GC phenotype was determined (D) and the number of NP-specific PNA<sup>+</sup>FAS<sup>+</sup> CD19<sup>+</sup> B cells was calculated (E). Data are representative of three independent experiments (mean  $\pm$  SD of five mice per group).

All p values were determined by two-tailed Student's t test. See also Figure S1.

that rIL-2 inhibited GC B cell development by a Treg cell-independent mechanism.

### Treatment of Mice with rIL-2 Impairs Tfh Cell Response

Tfh cells are required for the development and maintenance of GCs (Johnston et al., 2009; Nurieva et al., 2009; Yu et al., 2009). Because GCs are dependent on Tfh cell help, we next considered the possibility that the excess IL-2 compromised GC B cell formation by inhibiting

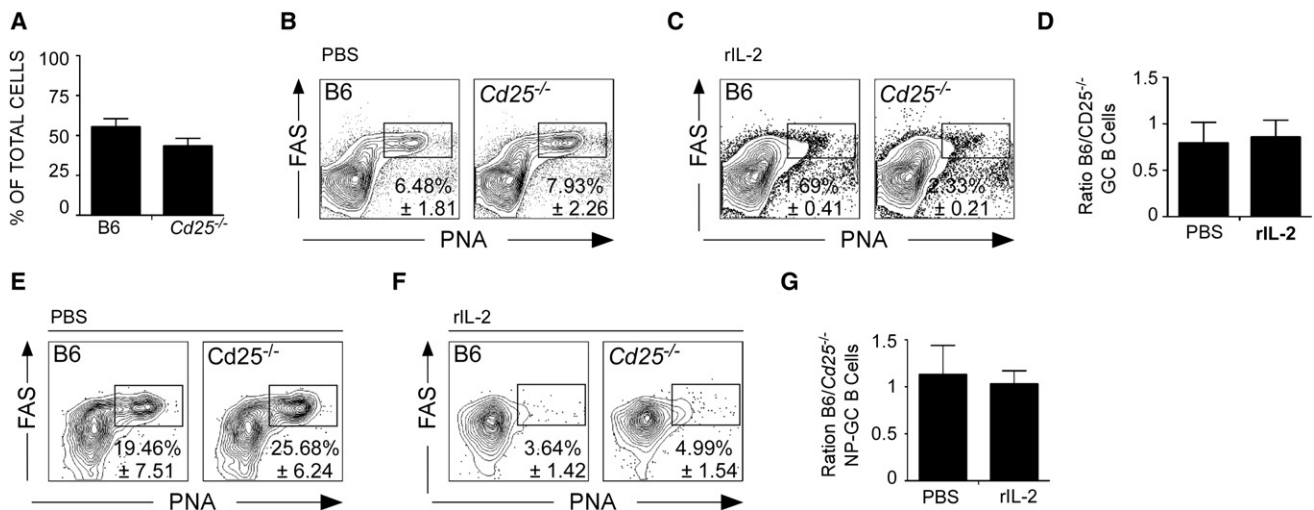
the development of Tfh cells. To test this possibility, we treated influenza-infected B6 mice with influenza, treated them with IL-2 or PBS, and enumerated Tfh cells in the mLN on day 10 post-infection. We characterized Tfh cells as PD-1<sup>hi</sup>CXCR5<sup>hi</sup> CD4<sup>+</sup> T cells (Figure 5A, left) that expressed high levels of the transcription factor Bcl6 and the costimulatory receptor ICOS compared to non-Tfh cells (Figure 5A, right). Importantly, we found fewer CD4<sup>+</sup> T cells that expressed Bcl6 after rIL-2 treatment (Figure 5B), and both the frequency (Figure 5C) and number (Figure 5D) of PD-1<sup>hi</sup>CXCR5<sup>hi</sup> Tfh cells were reduced relative to control mice.

To determine whether IL-2 was altering the maintenance of Tfh cells or preventing their development, we performed a similar experiment but examined the Tfh cell response on day 6, which is as early as we can consistently detect Tfh cells by flow cytometry. We found that IL-2 treatment between days 3 and 6 after infection reduced both the frequency (Figure 5E) and number (Figure 5F) of Tfh cells in the mLN. Thus, IL-2 acts early in the process of Tfh cell development and probably suppresses the commitment of primed CD4 T cells to the Tfh cell developmental pathway.

To test whether rIL-2 treatment disrupted Tfh cells by affecting the number of influenza-specific Tfh cells, we characterized the influenza nucleoprotein (NP)-specific CD4<sup>+</sup> T cell response by using a fluorochrome-labeled MHC class II tetramer (Figure 5G). We found that NP-specific CD4<sup>+</sup> T cells could be divided into

### rIL-2 Suppresses the GC B Cell Response to Influenza by a Treg Cell-Independent Mechanism

IL-2 signaling is crucial for the generation and maintenance of FoxP3-expressing Treg cells (Almeida et al., 2002; de la Rosa et al., 2004; Malek and Castro, 2010; Malek et al., 2002). Corresponding with this idea, we found more Treg cells in influenza-infected mice that were treated with rIL-2 than in control mice (Figure 4A). Because Treg cells suppress B cell immune responses, we next tested whether the inhibitory effect of rIL-2 treatment on GC formation was mediated by enhanced Treg cell function. Therefore, we treated FoxP3-DTR mice (Kim et al., 2007) with diphtheria toxin (DT), infected them with influenza virus, and treated them with either rIL-2 or PBS. Treatment of Foxp3-DTR mice with DT efficiently depleted FoxP3-expressing cells even after rIL-2 administration (Figure 4A). Infected mice were then administered rIL-2 on a daily basis, and we determined the frequency of PNA<sup>hi</sup>FAS<sup>+</sup> GC cells within CD19<sup>+</sup> B cells on day 10 after infection. As expected, FoxP3-DTR treated with rIL-2 alone had a decreased frequency (Figure 4B) and number (Figure 4C) of GC B cells compared to untreated FoxP3-DTR mice. Importantly, the frequency (Figure 4B) and number (Figure 4C) of GC B cells were even more reduced in Treg cell-depleted mice after rIL-2 treatment. As a control, DT administration to B6 mice did not affect the accumulation of GC B cell after influenza infection (data not shown). Thus, these results indicate



**Figure 3. rIL-2 Indirectly Inhibits GC B Cell Response to Influenza**

B6 mice were irradiated and reconstituted with a 50:50 mix of BM from wild-type CD45.1 donors and *Cd25*<sup>-/-</sup> donors (CD45.2).

(A) Percentage of leukocytes that expressed either CD45.1 or CD45.2 was determined in the mLN 8 weeks after reconstitution.

(B–D) Reconstituted mice were infected with PR8 and treated daily with 30,000 U of PBS (B) or rIL-2 (C) starting 3 days after infection, and the percentage of CD45.1<sup>+</sup> or CD45.2<sup>+</sup> CD19<sup>+</sup> B cells with a PNA<sup>+</sup>FAS<sup>+</sup> GC phenotype was determined on day 10. The ratio of B6 to *Cd25*<sup>-/-</sup> PNA<sup>+</sup>FAS<sup>+</sup> CD19<sup>+</sup> B cells was calculated (D).

(E–G) Reconstituted mice were infected with PR8 and treated daily with 30,000 U of rIL-2 or PBS starting 3 days after infection, and on day 10 the percentage of CD45.1<sup>+</sup> or CD45.2<sup>+</sup> NP-specific CD19<sup>+</sup> B cells with a PNA<sup>+</sup>FAS<sup>+</sup> GC phenotype was determined in the mLNs of mice treated with PBS (E) or with rIL-2 (F). The ratio of B6 to *Cd25*<sup>-/-</sup> PNA<sup>+</sup>FAS<sup>+</sup> NP-specific CD19<sup>+</sup> B cells was determined (G).

Data are representative of three independent experiments (mean ± SD of three to five mice per group). p values were determined by two-tailed Student's t test.

a PD-1<sup>hi</sup>CXCR5<sup>hi</sup> population, which expressed high levels of ICOS and Bcl6 (Tfh cells), and a PD-1<sup>lo</sup>CXCR5<sup>lo</sup> population, which expressed low levels of ICOS and Bcl6 (CD4<sup>+</sup> effectors) (Figure 5G). Interestingly, whereas NP-specific effector CD4<sup>+</sup> T cells were CD25<sup>hi</sup>, NP-specific Tfh cells were CD25<sup>lo</sup> (Figure 5G), suggesting that these populations received quantitatively different signals from the IL-2 receptor. We also found that IL-2 treatment reduced the frequency of NP-specific CD4<sup>+</sup> T cells that expressed Bcl6 (Figure 5H) and that the frequency (Figure 5I) and number (Figure 5J) of NP-specific Tfh cells were reduced in IL-2-treated mice compared to control. In contrast, rIL-2 treatment did not affect the accumulation of NP-specific effector CD4<sup>+</sup> T cells (Figure 5K). Given the reduced frequency of Bcl6-expressing Tfh cells in IL-2-treated mice, we next determined whether those Tfh cells remaining also expressed reduced amounts of Bcl6. We found that Bcl6 expression was identical in CD4<sup>+</sup>CXCR5<sup>+</sup>PD-1<sup>hi</sup> cells from IL-2-treated and control mice and in NP-specific CD4<sup>+</sup>CXCR5<sup>+</sup>PD-1<sup>hi</sup> cells from IL-2-treated and control mice (Figure 5L). Taken together, these results suggest that rIL-2 administration compromised the early commitment to the Tfh cell lineage rather than affecting already differentiated Tfh cells.

### IL-2 Directly Suppresses the Tfh Cell Response to Influenza

rIL-2 compromised the accumulation of Tfh cells, so we next considered the possibility that rIL-2 acted directly on CD4<sup>+</sup> T cells and inhibited Tfh cell development. To test this possibility, we generated mixed BM chimeras that contained a 50:50 mix of B6 and *Cd25*<sup>-/-</sup> cells. Two months after reconstitution, we infected

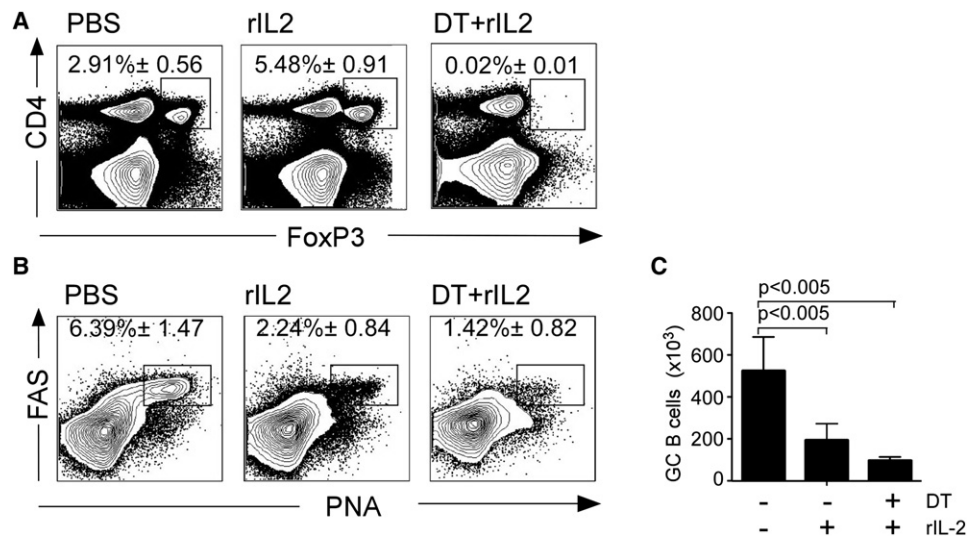
the chimeric mice with influenza, administered rIL-2 or PBS, and determined the frequency of Tfh cells within the B6 and *Cd25*<sup>-/-</sup> compartments on day 10 after infection. We found a higher frequency of Tfh cells in the *Cd25*<sup>-/-</sup> compartment than in the B6 compartment (Figure 6A). Similar results were obtained when we analyzed the frequency of NP-specific Tfh cells (Figure 6B). Thus, these results indicated that rIL-2 acts directly on T cells to inhibit Tfh cell development and that the loss of Tfh cells is the likely cause of poor GC and antibody responses in rIL-2-treated mice.

The previous results suggested that CD25 signaling negatively regulates Tfh cell development. If this was true, then we would predict a higher frequency of Tfh cells in the *Cd25*<sup>-/-</sup> CD4<sup>+</sup> T cell population compared to the B6 counterparts—even in untreated mice. To test this hypothesis, we infected B6:*Cd25*<sup>-/-</sup> mixed BM chimeras with influenza and calculated the frequency of Tfh cells within the B6 and *Cd25*<sup>-/-</sup> compartments 10 days postinfection. We found more *Cd25*<sup>-/-</sup> Tfh cells than B6 Tfh cells when we looked at total Tfh cells (Figure 6C) and NP-specific Tfh cells (Figure 6D). As a control, we also determined the relative frequencies of CD45.1 and CD45.2 cells in the total CD4<sup>+</sup> T cell compartment. We found that the frequencies of B6 and *Cd25*<sup>-/-</sup> T cells in the 50:50 chimeras were essentially identical (Figure 6E). Thus, our results indicated that lack of CD25 did not impair CD4<sup>+</sup> T cell development but did promote the differentiation of Tfh cells.

### DISCUSSION

Our data show that the availability of IL-2 regulates the differentiation of Tfh cells and thus controls GC formation and long-lived





**Figure 4. Inhibition of the GC B Cell Response after rIL-2 Treatment Does Not Require FoxP3<sup>+</sup> Treg Cells**

FoxP3-DTR mice were infected with PR8, administered PBS or DT on days 0, 4, and 7 after infection, or received DT on days 0, 4, and 7 after infection together with 30,000 U of human recombinant rIL-2 twice a day starting 3 days after infection. Cells from the mLN were analyzed on day 10.

(A and B) The percentage of CD4<sup>+</sup> T cells that expressed FoxP3 (A) and the percentage of CD19<sup>+</sup> B cells with a FAS<sup>+</sup>PNA<sup>+</sup> GC phenotype (B) were determined by flow cytometry.

(C) The number of FAS<sup>+</sup>PNA<sup>+</sup> GC B cells was calculated. Data are representative of two independent experiments (mean ± SD of five mice per time point). p values were determined by two-tailed Student's t test.

antibody responses. In conditions when excess IL-2 is available, Tfh cells are suppressed, GCs do not form, and influenza-specific IgG responses are reduced. Importantly, the lack of germinal center B cells in IL-2-treated mice is not a consequence of IL-2 signaling in B cells, but is the result of IL-2 signaling in T cells, which impairs Tfh cell differentiation. Moreover, although IL-2 signaling promotes the accumulation of Treg cells, which could suppress Tfh cells and GCs, Treg cells are not required for the ability of IL-2 to repress Tfh cell differentiation. Thus, our results demonstrate that IL-2 fine-tunes the balance of T effector cell and B cell helper aspects of the immune response to influenza.

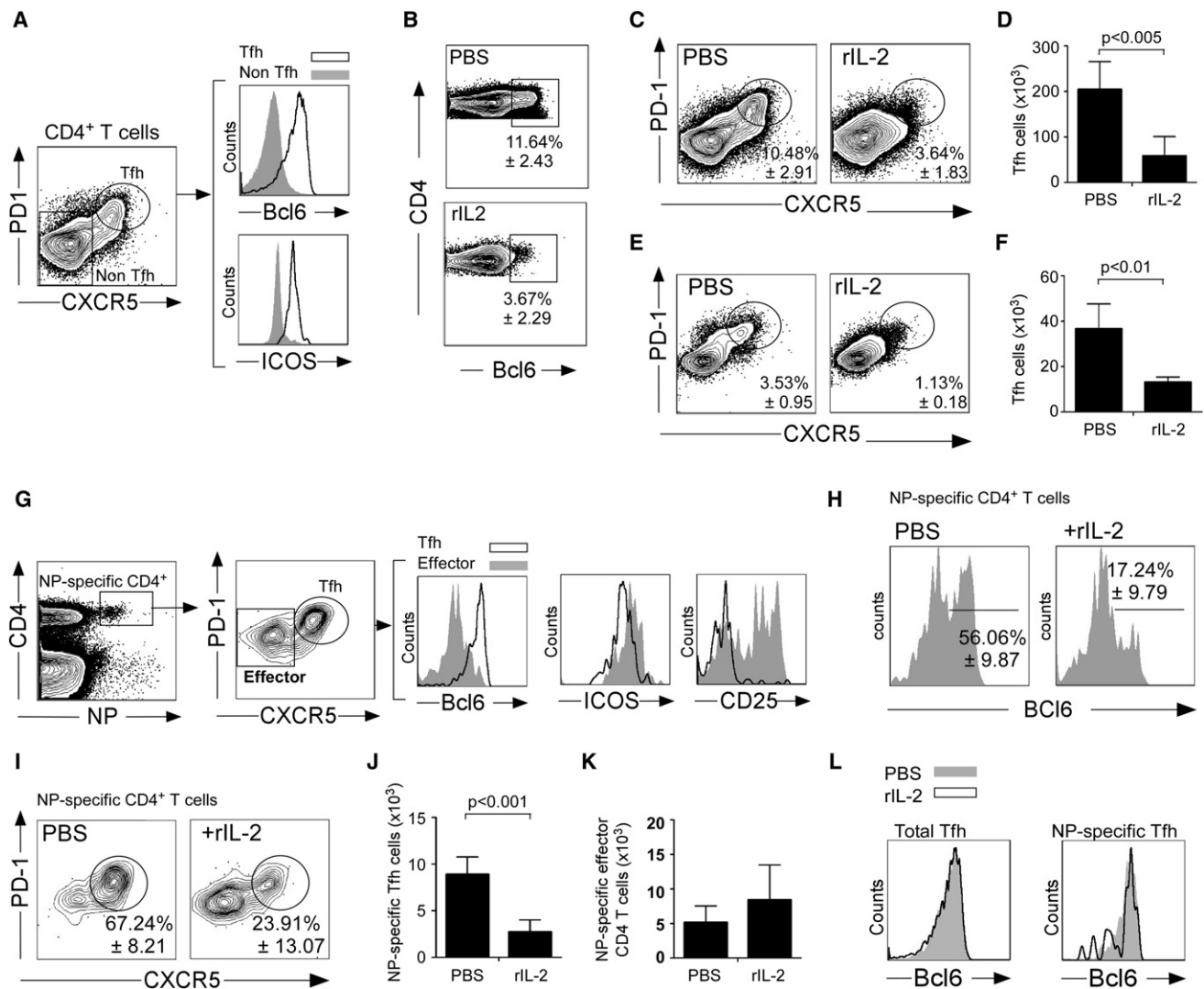
An important functional consequence of impaired Tfh cell differentiation is the reduced number of GC B cells and influenza-specific IgG-secreting cells. Although we cannot easily measure the affinity of the influenza-specific antibody produced in IL-2-treated mice, we anticipate that the antibodies that are produced would be of low affinity because of lack of selection in the GC. Moreover, in addition to reductions in long-lived, influenza-specific IgG-secreting cells in the BM, we would also anticipate that there would be defects in the generation of influenza-specific memory B cells, because both of these cell types are also thought to depend on events that occur in the GC. Thus, the ability of IL-2 to repress Tfh cell differentiation has multiple effects on the humoral immune response to influenza.

IL-2 is often perceived as a T cell growth factor for both CD4<sup>+</sup> and CD8<sup>+</sup> T cells (Malek, 2008). However, both CD4<sup>+</sup> and CD8<sup>+</sup> T cells respond to antigen and proliferate in mice lacking IL-2 or its receptor (Bachmann et al., 1995; Malek and Bayer, 2004; Williams et al., 2006). In fact, IL-2-deficient mice develop autoimmunity as a result of excessive effector T cell responses (Sadlack et al., 1993, 1995; Suzuki et al., 1995). This phenomenon can be

attributed, in part, to poor Treg cell survival in the absence of IL-2 signaling (de la Rosa et al., 2004; Malek et al., 2002). Although Treg cells clearly have the ability to impair CD4 T cell responses and even B cell responses (Alexander et al., 2011; Chung et al., 2011; Linterman et al., 2011), the role of IL-2 in preventing Tfh cell differentiation and GC formation does not appear to be dependent on Treg cells. Despite the presence of higher numbers of Treg cells in IL-2-treated mice, IL-2 still suppresses GC formation when Treg cells are eliminated. Moreover, only Cd25<sup>+/+</sup> Tfh cells are suppressed by the addition of IL-2 in B6:Cd25<sup>-/-</sup> chimeric mice, despite the equal opportunity of Treg cells to interact with both B6 and Cd25<sup>-/-</sup> T cells. Thus, IL-2 acts directly on differentiating Tfh precursor cells, rather than indirectly via Treg cell-mediated suppression.

The transcription factor Blimp1 controls the terminal differentiation of B cells into antibody-secreting plasma cells. As part of this pathway, Blimp1 represses the expression of Bcl6 (Johnston et al., 2009). Similarly, Blimp1 represses Bcl6 expression in T cells and promotes their terminal differentiation into effector cells (Crotty, 2011; Fazilleau et al., 2009; Poholek et al., 2010). Of course, impaired Bcl6 expression in T cells would also preclude the differentiation of Tfh cells (Johnston et al., 2009; Nurieva et al., 2009; Yu et al., 2009). Conversely, Bcl6 overexpression represses Blimp1 and favors Tfh cell development (Johnston et al., 2009). Thus, the balance between Blimp1 and Bcl6 expression is thought to control the relative commitment of CD4 T cells to effector or Tfh cell pathways (Crotty, 2011; Choi et al., 2011).

Interestingly, Blimp1 expression is increased in response to IL-2 (Gong and Malek, 2007). Thus, we speculate that one way in which IL-2 signaling impairs Tfh cell responses to influenza is by altering the Blimp1:Bcl6 ratio. This view is in agreement



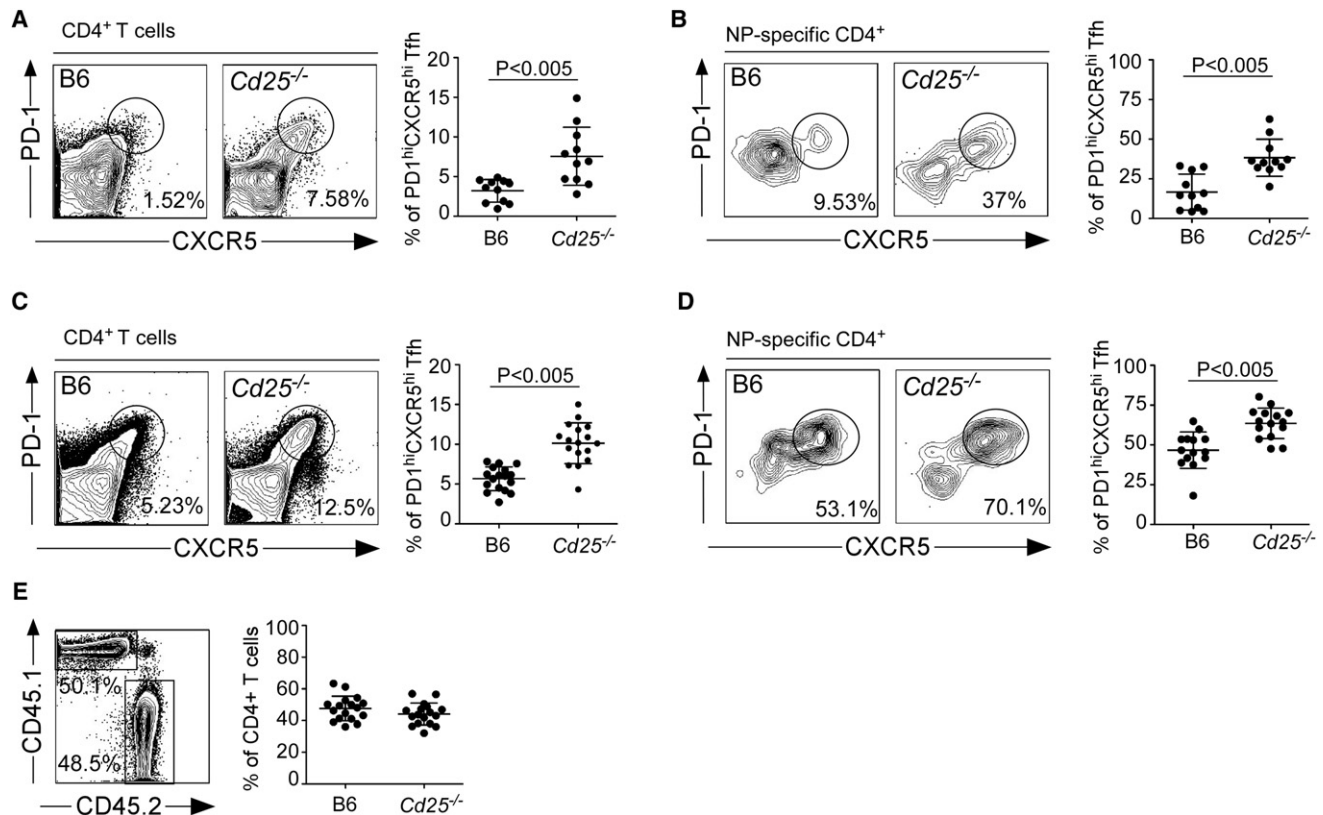
**Figure 5. rIL-2 Impairs the Tfh Cell Response to Influenza**

(A) B6 mice were infected with PR8 and the expression of Bcl6 and ICOS was evaluated on PD-1<sup>hi</sup>CXCR5<sup>hi</sup>CD4<sup>+</sup> T cells and PD-1<sup>lo</sup>CXCR5<sup>lo</sup>CD4<sup>+</sup> T cells on day 10. (B–D) B6 mice were infected with PR8, treated with 30,000 U of human rIL-2 or PBS twice a day starting 3 days after infection, and cells from the mLN were analyzed by flow cytometry on day 10. (B) The expression of Bcl6 in CD4<sup>+</sup> T cells was evaluated. (C) The percentage of CD4<sup>+</sup> T cells with a CXCR5<sup>hi</sup>PD-1<sup>hi</sup> Tfh cell phenotype was determined. (D) The number of CXCR5<sup>hi</sup>PD-1<sup>hi</sup> Tfh cells was calculated. (E and F) B6 mice were infected with PR8 and treated with 30,000 U of human rIL-2 or PBS twice a day starting 3 days after infection, and cells from the mLN were analyzed by flow cytometry on day 6. (E) The percentage of CD4<sup>+</sup> T cells with a CXCR5<sup>hi</sup>PD-1<sup>hi</sup> Tfh cell phenotype was determined. (F) The number of CXCR5<sup>hi</sup>PD-1<sup>hi</sup> Tfh cells was calculated. (G) Bcl6, ICOS, and CD25 expression was evaluated on PD-1<sup>hi</sup>CXCR5<sup>hi</sup> NP-specific CD4<sup>+</sup> T cells and PD-1<sup>lo</sup>CXCR5<sup>lo</sup> NP-specific CD4<sup>+</sup> T cells. (H) The expression of Bcl6 in NP-specific CD4<sup>+</sup> T cells was evaluated. (I) The percentage of NP-specific CD4<sup>+</sup> T cells with a CXCR5<sup>hi</sup>PD-1<sup>hi</sup> Tfh cell phenotype was determined. (J) The number of NP-specific CXCR5<sup>hi</sup>PD-1<sup>hi</sup> Tfh cells was calculated. (K) The number of NP-specific CXCR5<sup>lo</sup>PD-1<sup>lo</sup> effector CD4<sup>+</sup> T cells was calculated. (L) The expression of Bcl6 on CD4<sup>+</sup>CXCR5<sup>+</sup>PD-1<sup>hi</sup> cells and NP-specific CD4<sup>+</sup>CXCR5<sup>+</sup>PD-1<sup>hi</sup> cells from IL-2-treated and control mice was evaluated by flow cytometry.

Data are representative of four independent experiments (mean  $\pm$  SD of five mice per group).  $p$  values were determined by two-tailed Student's  $t$  test.

our results and previous results demonstrating that Blimp1 and Bcl6 are reciprocally expressed in CD25<sup>hi</sup> and CD25<sup>lo</sup> CD4<sup>+</sup> T cells responding to LCMV (Choi et al., 2011). Therefore, we

suggest that prolonged IL-2 signaling leads to increased Blimp1 expression and represses Bcl6 expression and thus impairs the differentiation or maintenance of Tfh cells. Interestingly, our data



**Figure 6. IL-2 Signaling Directly Inhibits Tfh Cell Responses to Influenza**

(A and B) B6 mice were irradiated and reconstituted with a 50:50 mix of BM from wild-type CD45.1 donors and *Cd25*<sup>-/-</sup> donors (CD45.2). Reconstituted mice were infected with PR8 and treated daily with 30,000 U of rIL-2 starting 3 days after infection, and cells from the mLN were analyzed on day 10. The percentage of CD45.1<sup>+</sup> or CD45.2<sup>+</sup> CD4<sup>+</sup>CXCR5<sup>+</sup>PD-1<sup>hi</sup> Tfh cells (A) and NP-specific CD45.1<sup>+</sup> or CD45.2<sup>+</sup> CD4<sup>+</sup>CXCR5<sup>+</sup>PD-1<sup>hi</sup> Tfh cells (B) was determined. Data were pooled from three independent experiments (mean  $\pm$  SD). Representative plots gated on CD4<sup>+</sup> T cells are shown.

(C and D) B6 mice were irradiated and reconstituted with a 50:50 mix of BM from wild-type CD45.1 donors and *Cd25*<sup>-/-</sup> donors (CD45.2). Reconstituted mice were infected with PR8, and cells from the mLN were analyzed on day 10. The percentage of CD45.1<sup>+</sup> or CD45.2<sup>+</sup> CD4<sup>+</sup>CXCR5<sup>+</sup>PD-1<sup>hi</sup> Tfh cells (C) and NP-specific CD45.1<sup>+</sup> or CD45.2<sup>+</sup> CD4<sup>+</sup>CXCR5<sup>+</sup>PD-1<sup>hi</sup> Tfh cells (D) was determined.

(E) The percentage of CD45.1<sup>+</sup> or CD45.2<sup>+</sup> total CD4<sup>+</sup> T cells was determined. Plot is gated on CD4<sup>+</sup> T cells. Data were pooled from three independent experiments (mean  $\pm$  SD). Representative plots gated on CD4<sup>+</sup> T cells are shown. p values were determined by two-tailed Student's t test.

show that although excess IL-2 suppresses Tfh cells, there is not a corresponding increase in the number of effector CD4<sup>+</sup> T cells. These results suggest that IL-2 signaling compromises the maintenance or survival of T cells committed to the Tfh cell pathway, rather than diverting influenza-specific CD4<sup>+</sup> T cells from the Tfh cell pathway into the Th1 effector cell pathway.

Published results show that IL-2 controls the differentiation of T cells by regulating the expression of a variety of transcription factors (Pipkin et al., 2010). For example, IL-2 represses ROR $\gamma$ t in CD4 T cells and prevents their differentiation into Th17 cells (Chen et al., 2011; Laurence et al., 2007; Pandiyan et al., 2011). IL-2 and STAT5 are also negative regulators of BCL6 in CD8<sup>+</sup> T cells and consequently, IL-2 signaling promotes the expansion of effector CD8<sup>+</sup> T cells at the expense of memory CD8<sup>+</sup> T cells (Kalia et al., 2010; Pipkin et al., 2010). Similarly, STAT5 inhibits the differentiation of Tfh cells via the BLIMP-1-dependent repression of BCL6 (Johnston et al., 2012). Consistent with this idea, we suggest that IL-2 signaling impairs or prevents Bcl6 expression in primed CD4<sup>+</sup> T cells, possibly via a STAT5-dependent mechanism (Johnston et al., 2012), and

prevents their subsequent differentiation into Tfh cells. In the absence of Bcl6, Tfh cell precursors do not express CXCR5, do not home to the T cell area, and do not receive the necessary signals from B cells to be maintained as Tfh cells (Choi et al., 2011; Deenick et al., 2011; Good-Jacobson et al., 2010). In contrast, once CD4<sup>+</sup> T cells have differentiated into Tfh cells, they lose CD25 expression and the addition of IL-2 does not affect Bcl6 or CXCR5 expression. Thus, we believe that IL-2 acts early in the Tfh cell differentiation pathway.

One of the hallmarks of Tfh cells is the expression of high levels of PD-1 (Yu and Vinuesa, 2010). PD-1 is an inhibitory receptor that is often associated with T cell exhaustion (Sharpe et al., 2007). However, PD-1 is also required for Tfh cell function and for germinal center formation (Good-Jacobson et al., 2010). However, it is not clear how PD-1 contributes to Tfh cell differentiation or maintenance. Interestingly, PD-1 signaling inhibits IL-2 production (Chikuma et al., 2009). Based on our observations showing that IL-2 signaling is deleterious to Tfh cell responses, we speculate that PD-1 signaling may promote the maintenance of Tfh cells by blocking autocrine IL-2 production.

Collectively, our results demonstrate that IL-2 plays a central role in the control of the Tfh cell response to influenza. Our findings also offer a new perspective for how Tfh cell development is regulated and demonstrate that the physiological availability of IL-2 is a critical factor that regulates successful B cell responses in vivo.

## EXPERIMENTAL PROCEDURES

### Mice, Infections, and In Vivo Treatments

C57BL/6 (B6), B6.129S4-*Il2ra*<sup>tm1Dw/J</sup>(*Cd25*<sup>-/-</sup>), and B6.*IgH<sup>g</sup>.Thy-1<sup>a</sup>.Ptrpc<sup>a</sup>* (CD45.1) mice were originally obtained from Jackson Laboratories and were bred in the University of Rochester animal facility. B6.129S6-*Foxp3*<sup>tm1DTR</sup> (FoxP3-DTR) mice were obtained from A. Rudensky (Memorial Sloan-Kettering Cancer Center). Influenza virus infections were performed intranasally with 500 egg infectious units of A/PR8/34 (PR8) in 100  $\mu$ l of PBS. *Heligmosomoides polygyrus* (Hp) infections were performed by gavage with 200 L3 Hp larvae as described (Wojciechowski et al., 2009). BM chimeric mice were generated by lethally irradiating CD45.1 mice with 950 Rads from a <sup>137</sup>Cs source delivered in two equal doses 4–5 hr apart. After irradiation, mice were intravenously injected with  $5 \times 10^6$  total BM cells and allowed to reconstitute for 8–12 weeks before influenza infection. In indicated experiments, WT and FoxP3-DTR mice received an intraperitoneal injection of 50  $\mu$ g/kg of DT (Sigma) on days 0, 4, and 7 after infection. For IL-2 treatment, mice were intraperitoneally administered 30,000 U of recombinant human IL-2 (National Cancer Institute) twice a day starting 3 days after infection. Control mice received injections of PBS. All experimental procedures involving animals were approved by the University of Rochester University Committee on Animal Resources and were performed according to guidelines outlined by the National Research Council.

### Cell Preparation and Flow Cytometry

Cell suspensions from mLNs were centrifuged, resuspended in 150 mM NH<sub>4</sub>Cl, 10 mM KHCO<sub>3</sub>, 0.1 mM EDTA for 10 min to lyse red cells, and filtered through a 70  $\mu$ m nylon cell strainer (BD Biosciences). Cells were washed and resuspended in PBS with 5% donor calf serum and 10  $\mu$ g/ml FcBlock (2.4G2; Truade Institute) for 10 min on ice followed by staining with fluorochrome-conjugated antibodies or tetramer reagents. The IA<sup>b</sup>NP<sub>311–325</sub> MHC class II tetramer was obtained from the NIH Tetramer Core Facility. Fluorochrome-labeled anti-CD45.1 (A20), anti-CD45.2 (104), anti-PD-1 (J43), anti-FoxP3 (FJK-16 s), anti-CD19 (1D3), anti-B220 (RA3-6B2), and anti-CD138 (281.2) were from eBioscience. Anti-Bcl6 (K112.91), anti-CXCR5 (2G-8), anti-ICOS (7E.17G9), anti-CD4 (RM4-5), and anti-CD95 (RM4-5) were from BD Biosciences. Peanut agglutinin (PNA) was obtained from Sigma (L7381; Sigma) and conjugated with FITC. Intracellular staining was performed with the mouse regulatory T cell staining kit (eBioscience) according to the manufacturer's instructions. Flow cytometry was performed with a FACSCanto II (BD Biosciences) or a C6 Flow Cytometer (Accuri) available through the Flow Cytometry Core Facility at the University of Rochester.

### Immunofluorescence

Lymph nodes were frozen in OCT (Tissue-Tek; Sakura) and 7  $\mu$ m frozen sections were prepared and stained as described (Rangel-Moreno et al., 2011). Slides were probed for 30 min at 25°C with anti-B220 (RA3-6B2), anti-IgD<sup>b</sup> (2170-170) obtained from BD Biosciences, anti-CD21 (7E9, BioLegend), and PNA (L7381; Sigma). Biotin-conjugated primary antibodies were detected with streptavidin-Alexa Fluor 555 (S21381; Invitrogen Life Sciences). Images were obtained with a Zeiss Axioplan 2 microscope with a Zeiss Axiocam digital camera (Zeiss) and saved as TIFF files.

### ELISAs and ELISPOT

For ELISAs, 96-well plates were coated with purified viral proteins (Lee et al., 2005) at 1  $\mu$ g/ml in 0.05 M Na<sub>2</sub>CO<sub>3</sub> (pH 9.6) overnight and then blocked for 1 hr with 1% BSA in PBS. Serum samples were serially diluted (3-fold) in PBS with 10 mg/ml BSA and 0.1% Tween 20 before incubation on coated plates. After washing, bound antibody was detected with HRP-conjugated goat anti-mouse IgG1 (Southern Biotechnology Associates). For ELISPOT, multiscreen

cellulose filter plates (Millipore) were coated overnight with 1  $\mu$ g/ml purified viral proteins. Cells were collected from bone marrow and plated in triplicate starting at  $1 \times 10^6$ /well and in 3-fold serial dilutions in complete RPMI 1640 containing 10% FBS. After 5 hr, the wells were washed with PBS containing 0.5% BSA and 0.05% Tween 20, and IgG was detected with alkaline phosphatase-conjugated goat anti-mouse IgG (Southern Biotechnology Associates). Spots were automatically counted in a CTL-immunoSpot S5 analyzer.

### NP-Tetramer Preparation for Detection of Influenza-Specific B Cells

The coding sequence of NP from the A/PR8/34 virus was synthesized in-frame with the coding sequence for a 15 amino acid biotinylation consensus site (Beckett et al., 1999) on the 3' end (GeneArt, Regensburg, Germany). The modified NP sequence was cloned in-frame to the 6 $\times$  His tag in the pTRC-His2c expression vector (Invitrogen, Carlsbad, CA). NP protein was expressed in *E. coli* strain Ultra BL21 (DE3) (EdgeBio, Gaithersburg, MD) and purified by FPLC with a HisTrap HP Column (GE Healthcare). Purified protein was biotinylated by addition of Biotin-protein ligase (Avidity, Aurora, CO). The biotinylated NP protein was then tetramerized with PE-labeled streptavidin (Prozyme, Hayward, CA). Labeled tetramers were purified by size exclusion on a HiPrep 16/60 Sephacryl S-300 column (GE Healthcare, Piscataway, NJ).

### Statistical Analysis

The statistical significance of differences in mean values was analyzed by a two-tailed Student's t test. p values of less than 0.05 were considered statistically significant.

### SUPPLEMENTAL INFORMATION

Supplemental Information includes one figure and can be found with this article online at doi:10.1016/j.immuni.2012.02.012.

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